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PCT09
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/856,050

DATE: 10/11/2001
TIME: 09:50:56

Input Set : A:\sequence listing.txt
Output Set: N:\CRF3\10112001\I856050.raw

3 <110> APPLICANT: UEMURA, Hidetoshi
4 OKUI, Akira
5 KOMINAMI, Katsuya
6 YAMAGUCHI, Nozomi
7 MITSUI, Shinichi
9 <120> TITLE OF INVENTION: PROTEIN EXPRESSION VECTOR AND USE THEREOF
11 <130> FILE REFERENCE: UEMURA=8
13 <140> CURRENT APPLICATION NUMBER: 09/856,050
14 <141> CURRENT FILING DATE: 2001-05-17
16 <150> PRIOR APPLICATION NUMBER: JP 10/331515
17 <151> PRIOR FILING DATE: 1998-11-20
19 <150> PRIOR APPLICATION NUMBER: PCT/JP99/06474
20 <151> PRIOR FILING DATE: 1999-11-19
22 <160> NUMBER OF SEQ ID NOS: 22
24 <170> SOFTWARE: PatentIn version 3.1
26 <210> SEQ ID NO: 1
27 <211> LENGTH: 117
28 <212> TYPE: DNA
29 <213> ORGANISM: Artificial Sequence
31 <220> FEATURE:
32 <223> OTHER INFORMATION: Synthetic ✓
34 <220> FEATURE:
35 <221> NAME/KEY: misc_feature
36 <223> OTHER INFORMATION: Designed oligonucleotide to construct plasmid pTrypHis
39 <400> SEQUENCE: 1
40 aagcttggct agcaacacca tgaatctact cctgatcctt acctttgttg ctgctgctgt 60
42 tgctgcccccc tttcaccatc accatcacca tgacgacgat gacaaggatc cgaattc 117
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47 <212> TYPE: DNA
48 <213> ORGANISM: Artificial Sequence
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53 <220> FEATURE:
54 <221> NAME/KEY: misc_feature
55 <223> OTHER INFORMATION: Designed oligonucleotide to construct plasmid pTrypHis
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59 gaattcggat ccttgcatac gtcgtcatgg tcatggat ggtgaaagg ggcagcaaca 60
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67 <213> ORGANISM: Artificial Sequence
69 <220> FEATURE:
70 <223> OTHER INFORMATION: Synthetic ✓
72 <220> FEATURE:
73 <221> NAME/KEY: misc_feature

ENTERED

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74 <223> OTHER INFORMATION: Designed oligonucleotide primer to amplify neurosin-encoding
 sequ 75 ence

78 <400> SEQUENCE: 3

79 ttgggtgcattt gcggtt

82 <210> SEQ ID NO: 4

83 <211> LENGTH: 20

84 <212> TYPE: DNA

85 <213> ORGANISM: Artificial Sequence

87 <220> FEATURE:

88 <223> OTHER INFORMATION: Synthetic ✓

90 <220> FEATURE:

91 <221> NAME/KEY: misc_feature

92 <223> OTHER INFORMATION: Designed oligonucleotide primer to amplify neurosin-encoding
 sequ 93 ence

96 <400> SEQUENCE: 4

97 ggaatttactt tggcctgaaat

100 <210> SEQ ID NO: 5

101 <211> LENGTH: 26

102 <212> TYPE: DNA

103 <213> ORGANISM: Artificial Sequence

105 <220> FEATURE:

106 <223> OTHER INFORMATION: Synthetic ✓

108 <220> FEATURE:

109 <221> NAME/KEY: misc_feature

110 <223> OTHER INFORMATION: Designed oligonucleotide primer to amplify a portion of
 plasmid p

111 TrypHis/Neurosin

114 <400> SEQUENCE: 5

115 ctaagcttga cggacgtatgtac aaggttt

118 <210> SEQ ID NO: 6

119 <211> LENGTH: 27

120 <212> TYPE: DNA

121 <213> ORGANISM: Artificial Sequence

123 <220> FEATURE:

124 <223> OTHER INFORMATION: Synthetic ✓

126 <400> SEQUENCE: 6

127 tcctcgagac ttggcctgaa tggtttt

130 <210> SEQ ID NO: 7

131 <211> LENGTH: 26

132 <212> TYPE: DNA

133 <213> ORGANISM: Artificial Sequence

135 <220> FEATURE:

136 <223> OTHER INFORMATION: Synthetic ✓

138 <220> FEATURE:

139 <221> NAME/KEY: misc_feature

140 <223> OTHER INFORMATION: Designed oligonucleotide primer to amplify a portion of
 plasmid p

141 TrypHis/Neurosin

144 <400> SEQUENCE: 7

145 ccaagcttca ccatcaccat caccat

148 <210> SEQ ID NO: 8

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Input Set : A:\sequence listing.txt

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149 <211> LENGTH: 99
 150 <212> TYPE: DNA
 151 <213> ORGANISM: Artificial Sequence
 153 <220> FEATURE:
 154 <223> OTHER INFORMATION: Synthetic O^V
 156 <220> FEATURE:
 157 <221> NAME/KEY: misc_feature
 158 <223> OTHER INFORMATION: Designed oligonucleotide to construct plasmid pSecTrypHis
 161 <400> SEQUENCE: 8
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 164 tgctgcccccc tttgacgacg atgacaaggaa tccgaattc 99
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 170 <213> ORGANISM: Artificial Sequence
 172 <220> FEATURE:
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 175 <220> FEATURE:
 176 <221> NAME/KEY: misc_feature
 177 <223> OTHER INFORMATION: Designed oligonucleotide to construct plasmid pSecTrypHis
 180 <400> SEQUENCE: 9
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 189 <213> ORGANISM: Artificial Sequence
 191 <220> FEATURE:
 192 <223> OTHER INFORMATION: Synthetic O^V
 194 <220> FEATURE:
 195 <221> NAME/KEY: misc_feature
 196 <223> OTHER INFORMATION: Designed oligonucleotide primer to amplify a portion of plasmid p
 197 SecTrypHis/Neurosin
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 205 <211> LENGTH: 29
 206 <212> TYPE: DNA
 207 <213> ORGANISM: Artificial Sequence
 209 <220> FEATURE:
 210 <223> OTHER INFORMATION: Synthetic O^V
 212 <220> FEATURE:
 213 <221> NAME/KEY: misc_feature
 214 <223> OTHER INFORMATION: Designed oligonucleotide primer to amplify a portion of plasmid p
 215 SecTrypHis/Neurosin
 218 <400> SEQUENCE: 11
 219 tgaagcttgc catggaccaa cttgtcatc 29
 222 <210> SEQ ID NO: 12
 223 <211> LENGTH: 17

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Input Set : A:\sequence listing.txt
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224 <212> TYPE: DNA
 225 <213> ORGANISM: Artificial Sequence
 227 <220> FEATURE:
 228 <223> OTHER INFORMATION: Synthetic *OV*
 230 <220> FEATURE:
 231 <221> NAME/KEY: misc_feature
 232 <223> OTHER INFORMATION: Designed oligonucleotide primer to amplify a portion of
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 233 TrypSigTag
 236 <400> SEQUENCE: 12
 237 gcacagtcga ggctgat 17
 240 <210> SEQ ID NO: 13
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 245 <220> FEATURE:
 246 <223> OTHER INFORMATION: Synthetic *OV*
 248 <220> FEATURE:
 249 <221> NAME/KEY: misc_feature
 250 <223> OTHER INFORMATION: Designed oligonucleotide primer to amplify a portion of
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 251 FBTrypSigTag
 254 <400> SEQUENCE: 13
 255 caaatgtggc atggctg 17
 258 <210> SEQ ID NO: 14
 259 <211> LENGTH: 672
 260 <212> TYPE: DNA
 261 <213> ORGANISM: Homo sapiens
 263 <220> FEATURE:
 264 <221> NAME/KEY: CDS
 265 <222> LOCATION: (1)..(672)
 266 <223> OTHER INFORMATION:
 269 <400> SEQUENCE: 14
 270 ttg gtg cat ggc gga ccc tgc gac aag aca tct cac ccc tac caa gct 48
 271 Leu Val His Gly Gly Pro Cys Asp Lys Thr Ser His Pro Tyr Gln Ala
 272 1 5 10 15
 274 gcc ctc tac acc tcg ggc cac ttg ctc tgt ggt ggg gtc ctt atc cat 96
 275 Ala Leu Tyr Thr Ser Gly His Leu Leu Cys Gly Val Leu Ile His
 276 20 25 30
 278 cca ctg tgg gtc ctc aca gct gcc cac tgc aaa aaa ccg aat ctt cag 144
 279 Pro Leu Trp Val Leu Thr Ala Ala His Cys Lys Lys Pro Asn Leu Gln
 280 35 40 45
 282 gtc ttc ctg ggg aag cat aac ctt cgg caa agg gag agt tcc cag gag 192
 283 Val Phe Leu Gly Lys His Asn Leu Arg Gln Arg Glu Ser Ser Gln Glu
 284 50 55 60
 286 cag agt tct gtt gtc cgg gct gtg atc cac cct gac tat gat gcc gcc 240
 287 Gln Ser Ser Val Val Arg Ala Val Ile His Pro Asp Tyr Asp Ala Ala
 288 65 70 75 80
 290 agc cat gac cag gac atc atg ctg ttg cgc ctg gca cgc cca gcc aaa 288
 291 Ser His Asp Gln Asp Ile Met Leu Leu Arg Leu Ala Arg Pro Ala Lys
 292 85 90 95

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| | | | | | | | | | | | | | | | | | |
|-----|---|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| 294 | ctc | tct | gaa | ctc | atc | cag | ccc | ctt | ccc | ctg | gag | agg | gac | tgc | tca | gcc | 336 |
| 295 | Leu | Ser | Glu | Leu | Ile | Gln | Pro | Leu | Pro | Leu | Glu | Arg | Asp | Cys | Ser | Ala | |
| 296 | | | | 100 | | | | | 105 | | | | | 110 | | | |
| 298 | aac | acc | acc | agc | tgc | cac | atc | ctg | ggc | tgg | ggc | aag | aca | gca | gat | ggt | 384 |
| 299 | Asn | Thr | Thr | Ser | Cys | His | Ile | Leu | Gly | Trp | Gly | Lys | Thr | Ala | Asp | Gly | |
| 300 | | | | 115 | | | | 120 | | | | 125 | | | | | |
| 302 | gat | ttc | cct | gac | acc | atc | cag | tgt | gca | tac | atc | cac | ctg | gtg | tcc | cgt | 432 |
| 303 | Asp | Phe | Pro | Asp | Thr | Ile | Gln | Cys | Ala | Tyr | Ile | His | Leu | Val | Ser | Arg | |
| 304 | | 130 | | | | 135 | | | | 140 | | | | | | | |
| 306 | gag | gag | tgt | gag | cat | gcc | tac | cct | ggc | cag | atc | acc | cag | aac | atg | ttg | 480 |
| 307 | Glu | Glu | Cys | Glu | His | Ala | Tyr | Pro | Gly | Gln | Ile | Thr | Gln | Asn | Met | Leu | |
| 308 | 145 | | | 150 | | | | | 155 | | | 160 | | | | | |
| 310 | tgt | gct | ggg | gat | gag | aag | tac | ggg | aag | gat | tcc | tgc | cag | ggt | gat | tct | 528 |
| 311 | Cys | Ala | Gly | Asp | Glu | Lys | Tyr | Gly | Lys | Asp | Ser | Cys | Gln | Gly | Asp | Ser | |
| 312 | | | | 165 | | | | 170 | | | 175 | | | | | | |
| 314 | ggg | ggt | ccg | ctg | gta | tgt | gga | gac | cac | ctc | cga | ggc | ctt | gtg | tca | tgg | 576 |
| 315 | Gly | Gly | Pro | Leu | Val | Cys | Gly | Asp | His | Leu | Arg | Gly | Leu | Val | Ser | Trp | |
| 316 | | | | 180 | | | | 185 | | | 190 | | | | | | |
| 318 | ggt | aat | atc | ccc | tgt | gga | tca | aag | gag | aag | cca | gga | gtc | tac | acc | aac | 624 |
| 319 | Gly | Asn | Ile | Pro | Cys | Gly | Ser | Lys | Glu | Lys | Pro | Gly | Val | Tyr | Thr | Asn | |
| 320 | | | | 195 | | | | 200 | | | 205 | | | | | | |
| 322 | gtc | tgc | aga | tac | acg | aac | tgg | atc | caa | aaa | acc | att | cag | gcc | aag | tga | 672 |
| 323 | Val | Cys | Arg | Tyr | Thr | Asn | Trp | Ile | Gln | Lys | Thr | Ile | Gln | Ala | Lys | | |
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| 327 | <210> SEQ ID NO: 15 | | | | | | | | | | | | | | | | |
| 328 | <211> LENGTH: 223 | | | | | | | | | | | | | | | | |
| 329 | <212> TYPE: PRT | | | | | | | | | | | | | | | | |
| 330 | <213> ORGANISM: Homo sapiens | | | | | | | | | | | | | | | | |
| 332 | <220> FEATURE: | | | | | | | | | | | | | | | | |
| 333 | <223> OTHER INFORMATION: Synthetic 0 | | | | | | | | | | | | | | | | |
| 335 | <400> SEQUENCE: 15 | | | | | | | | | | | | | | | | |
| 337 | Leu | Val | His | Gly | Gly | Pro | Cys | Asp | Lys | Thr | Ser | His | Pro | Tyr | Gln | Ala | |
| 338 | 1 | | | | 5 | | | | 10 | | | | 15 | | | | |
| 341 | Ala | Leu | Tyr | Thr | Ser | Gly | His | Leu | Leu | Cys | Gly | Val | Leu | Ile | His | | |
| 342 | | | | 20 | | | | | 25 | | | 30 | | | | | |
| 345 | Pro | Leu | Trp | Val | Leu | Thr | Ala | Ala | His | Cys | Lys | Lys | Pro | Asn | Leu | Gln | |
| 346 | | | | 35 | | | | 40 | | | 45 | | | | | | |
| 349 | Val | Phe | Leu | Gly | Lys | His | Asn | Leu | Arg | Gln | Arg | Glu | Ser | Ser | Gln | Glu | |
| 350 | | | | 50 | | | | 55 | | | 60 | | | | | | |
| 353 | Gln | Ser | Ser | Val | Val | Arg | Ala | Val | Ile | His | Pro | Asp | Tyr | Asp | Ala | Ala | |
| 354 | 65 | | | | 70 | | | | 75 | | | 80 | | | | | |
| 357 | Ser | His | Asp | Gln | Asp | Ile | Met | Leu | Leu | Arg | Leu | Ala | Arg | Pro | Ala | Lys | |
| 358 | | | | | 85 | | | | 90 | | | 95 | | | | | |
| 361 | Leu | Ser | Glu | Leu | Ile | Gln | Pro | Leu | Pro | Leu | Glu | Arg | Asp | Cys | Ser | Ala | |
| 362 | | | | | 100 | | | | 105 | | | 110 | | | | | |
| 365 | Asn | Thr | Thr | Ser | Cys | His | Ile | Leu | Gly | Trp | Gly | Lys | Thr | Ala | Asp | Gly | |
| 366 | | | | 115 | | | | 120 | | | 125 | | | | | | |
| 369 | Asp | Phe | Pro | Asp | Thr | Ile | Gln | Cys | Ala | Tyr | Ile | His | Leu | Val | Ser | Arg | |
| 370 | | | | 130 | | | | 135 | | | 140 | | | | | | |

VERIFICATION SUMMARY

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Input Set : A:\sequence listing.txt

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